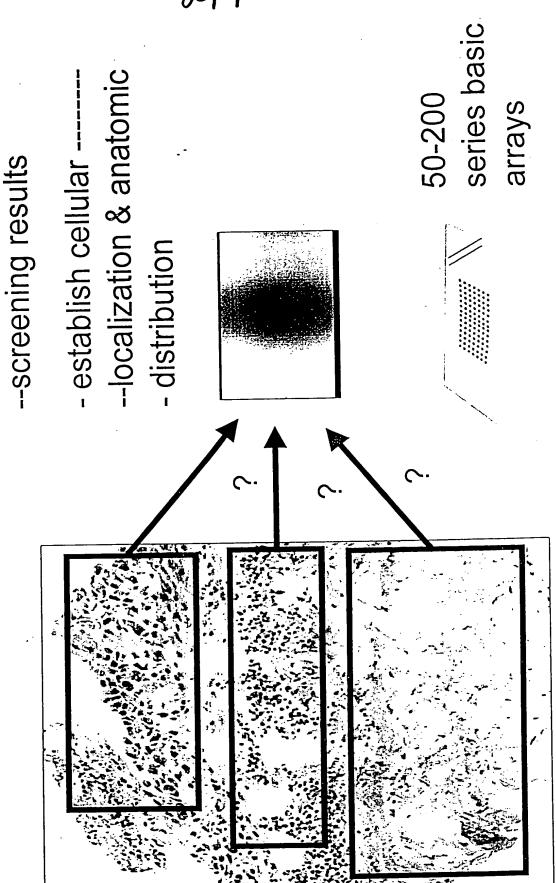
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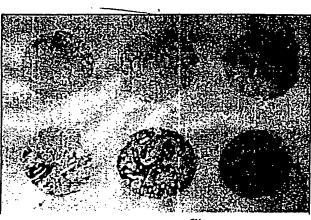
F16. 1A

- verification of initial

### Validation - phase



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FIG. 2A		F16. 28



Six tissue elements from a PathWork<sup>TM</sup> Breast Array BR50 stained with CK7 showing selective staining (400X). Tissue elements are 0.6mm in diameter and spaced 0.3mm apart.

F16. 2C

Sample Data Sheet Information (for one tissue element, located at A1a) provided with every pathWorks<sup>™</sup> Tissue MicroArray

Cat. #	BR50
Lot#	07
Coordinates:	A1a
Organ:	Breast
Category:	Breast cancer, ductal
Sex:	F
Age:	67
BRE Grade:	G2
Tubuli:	3
Polymorphy:	3

Mitoses:	1
PT:	1
pN:	0
LN pos	0
LN all	23
Tumor dia.	15
Surgery:	mastectomy

#16. 2D

FIG 2E

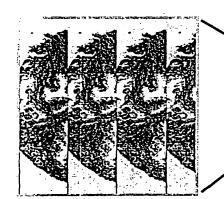
## Tissue MicroArrays



High-throughput in situ analysis of DNA, RNA and protein

Discovery of diagnostic and prognostic correllations

Target prioritization



# The Clinomics Biorepository

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000007489 Breast cancer	IPA, A, tubes	SIF	1100 mg/l			
000007491 Breast cancer	TPA, B, bags	- JFFPE	₹ 100 Mg	0		
000007492 Breast cancer	TPA, B, bags	- JFFPE	100 mg	0		74
000007493 Breast cancer	TPA, A, tubes	FFPE	100 mg	0		
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FIG. 4A

## **Biorepository Database**

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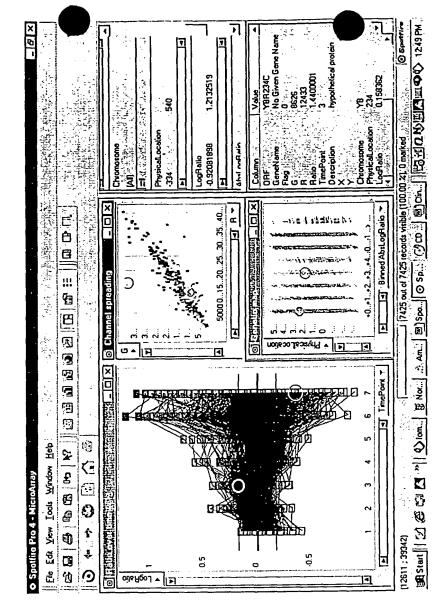
Medication and treatment histories, with outcomes, and expression profiles

## **Biorepository Database**

Sorting and selection of patient populations

Clustering analysis

expression profiles vs. Development of medical data



Clustering analysis